

~~No. 08/710,633, which issued on February 6, 2001 as US Patent~~  
No. 6,184,344, and which was a national stage application under  
35 U.S.C. § 371 of International Application No. PCT/US95/05668,  
filed May 4, 1995, which International Application was published in  
~~English~~

Government Rights:

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The invention disclosed herein was supported in part  
by Grants Number R01 GM 48897 and P01 GM 48870 from  
the National Institutes of Health. The United States  
government may have certain rights to this invention.

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Page 1, lines 13-18, please delete the statement of Government Rights.

Page 7, first paragraph, lines 2-19, please substitute the following replacement  
paragraph:

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One aspect of the invention is directed to a method of native  
chemical ligation. The method of native chemical ligation facilitates the  
chemical synthesis of proteins and large oligopeptide. The principle of  
'native chemical ligation' is shown in FIG. 1. The first step is the  
chemoselective reaction of an unprotected synthetic peptide- $\alpha$ - thioester  
with another unprotected peptide segment containing an N-terminal Cys  
residue, to give a thioester-linked intermediate as the initial covalent  
product. Without change in the reaction conditions, this intermediate  
undergoes spontaneous, rapid intramolecular reaction to form a native  
peptide bond at the ligation site. The target full length polypeptide product  
is obtained in the desired final form without further manipulation. The  
general synthetic access provided by the method of native chemical

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ligation greatly expands the scope of variation of the covalent structure of the protein molecule.

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Page 9, last paragraph, line 28 to line 6 of page 10, please substitute the following replacement paragraph:

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The oligopeptide thioester ( $\alpha$ -COSR moiety) of FIG. 1 can be readily generated from a corresponding oligopeptide thiol ( $\alpha$ COSH) prepared by highly optimized stepwise SPPS on a thioester resin. The thioester resin was prepared by the method of L. E. Canne et al., Tetrahedron Letters (1995): vol. 36, pp. 1217-1220, incorporated herein by reference. The method of Canne employs the thioester resin disclosed by Blake and Yamashiro (J. Blake, Int. J. Pept. Protein Res. (1981): vol. 17, p 273; D. Yamashiro, et al., Int. J. Pept. Protein Res. (1988): vol. 31, p 322). Peptide products were cleaved, purified, and characterized by conventional methods. (M. Schnolzer, et al., Int. J. Pept. Protein Res., (1992): vol. 40, pp 180-193.)

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Page 10, last paragraph, line 29 to line 14 of page 11, please substitute the following replacement paragraph:

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The invention stated herein does not require the use of any protecting groups for the coupling of two oligopeptides because a less reactive (and thus more chemoselective) thioester electrophile is used instead of the acyl disulfide moiety (Yamashiro's approach). In the intermolecular coupling step, this thioester electrophile requires a more nucleophilic sulfhydryl moiety rather than a free amine. The nucleophilic sulfhydryl moiety can be found on cysteine residues. Since the amino and

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hydroxyl functionalities are relatively unreactive to the thioester electrophile, a selective coupling of the two unprotected oligopeptides is achieved with the cysteine sulfhydryl moiety. The sulfhydryl group on the cysteine of peptide 2 will first attack the thioester of peptide 1 and form a coupled thioester intermediate. This coupled thioester intermediate is concomitantly attacked by the free  $\alpha$ -amino moiety from the cysteine and spontaneously rearranges to form the native peptide bond. Yields are therefore increased by eliminating protection and deprotection steps, since side undesired reactions are reduced (FIG. 1).

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Page 11, last paragraph, line 35 to line 9 of page 12, please substitute the following replacement paragraph:

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In this method (FIG. 2), thiol 3 is generated from the reaction of chloride 2 (Yamashiro et. al. Int. J. Pept. Protein Res. (1988): vol. 31, p 322) with thiourea, followed by hydrolysis of the resulting thiuronium salt in aqueous base. Thiol 3 is a general intermediate which can be reacted with a wide range of commercially available Boc-amino acid succinimide esters to produce the desired thioester linker 1 which is conveniently isolated as the dicyclohexylamine (DCHA) salt.

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Page 12, please delete the second paragraph starting on line 10 and extending to line 9 on page 14 and substitute the following amended paragraph:

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Model studies were undertaken with small peptides to investigate the native chemical ligation approach. To help explore the mechanism of the reaction, the peptide Leu-Tyr-Arg-Ala-Gly- $\alpha$ -COSBzl (SEQ ID NO 3) was reacted with Ac-Cys. The exact mass of the resulting ligation product

was determined by electrospray mass spectrometry, and was consistent with a thioester-linked peptide as the ligation product generated by nucleophilic attack of the Ac-Cys side chain on the  $\alpha$ -thioester moiety of the peptide. Reaction of Leu-Tyr-Arg-Ala-Gly- $\alpha$ -COSBzl (SEQ ID NO 3) with H-Cys-Arg-Ala-Glu-Tyr-Ser (SEQ ID NO 2) (containing an unblocked  $\alpha$ -NH<sub>2</sub> functional group) proceeded rapidly at pH 6.8 (below pH 6 the reaction proceeded very slowly, suggesting the involvement of the ionized thiolate form of the Cys side chain), and gave a single product of the expected mass. This product lacked susceptibility to nucleophiles, and had the ability to form disulfide-linked dimeric peptides, indicating unambiguously the formation of a native amide bond at the ligation site. These studies were consistent with the mechanism shown in FIG 1, in which the initial thioester ligation product was not observed as a discrete intermediate because of the rapid rearrangement to form a stable peptide bond. Facile intramolecular reaction results from the favorable geometric arrangement of the  $\alpha$ -NH<sub>2</sub> moiety with respect to the thioester formed in the initial chemoselective ligation reaction. Use of such 'entropy activation' for peptide bond formation is based on principles enunciated by Brenner. (M. Brenner, in Peptides. Proceedings of the Eighth European Peptide Symposium H. C. Beyerman, Eds. (North Holland, Amsterdam, 1967) pp. 1-7.) The concept of 'entropy activation' for peptide bond formation has been more recently adopted by D. S. Kemp et al. (J. Org. Chem. (1993): vol. 58, p 2216) and by C.-F. Liu, et al. (J. Am. Chem. Soc. (1994): vol. 116, p 4149).

Page 16, please delete the second paragraph starting on line 3 and extending to line 18 on page 14 and substitute the following amended paragraph:

B7 Further model reactions demonstrate that the use of better thioester leaving groups results in faster ligation reactions. We applied this observation to the native chemical ligation of peptides from the extracellular domain of a human cytokine receptor (R. D'Andrea, et al., Blood, (1994): vol. 83, p 2802.) as shown in FIG. 5. Use of the 5-thio-2-nitrobenzoic acid (-SNB) leaving group, corresponding to the reduced form of Elman's reagent, gave rapid high yield reaction. As described below in connection with FIG. 5, the reaction between the peptide segments was observed to have gone essentially to completion in less than 5 minutes, giving the 50 residue product with a native peptide bond at the site of ligation. Thus, rapid native chemical ligation can be achieved by use of a thioester leaving group with suitably tuned properties.

Page 16, please delete the last paragraph starting on line 19 and extending to line 18 on page 18 and substitute the following amended paragraph:

B8 Application of the native chemical ligation method to the total synthesis of a protein molecule was illustrated by the preparation of human interleukin 8 (IL-8). (M. Baggiolini, et al., FEBS Lett. (1989): vol. 307, p 97; I. Clark-Lewis, et al., J. Biol. Chem. (1994): vol. 269, p 16075 (1994); I. Clark-Lewis, Biochemistry (1991): vol. 30, p 3128; and K. Rajarathnam, et al., (1994): Biochemistry, (1994): vol. 29, p 1689.) The 72 amino acid polypeptide chain contains four Cys residues, which form two functionally critical disulfide bridges in the native protein molecule. The total synthesis of IL-8 is shown in FIG. 7. The two unprotected synthetic peptide segments reacted cleanly to give the full length polypeptide chain in reduced form without further chemical manipulation (9). This successful

ligation was particularly significant because the 33- and 39-residue IL-8 segments each contained two Cys residues, and together encompassed 18 of the 20 genetically encoded amino acids found in proteins. The purified product was folded and oxidized as previously described, to give IL-8 with a mass precisely 4 daltons less than that of the original ligation product, indicating the formation of two disulfide bonds. The properties of this folded product were identical to those of previously studied authentic IL-8 samples. Titration in an assay for neutrophil elastase release demonstrated that the potencies ( $ED_{50}=0.3nM$ ) and maximal responses of the folded, ligated [Ala33]IL-8 and the corresponding molecule obtained by conventional synthesis were indistinguishable and identical to native sequence IL-8. This result unambiguously confirmed the formation of a peptide bond at the ligation site, because the thioester-to-amide rearrangement must have taken place to give the free Cys<sup>34</sup> side chain that formed the native disulfide bond (see FIG. 7).

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Page 20, please delete the second paragraph, lines 6-12, and substitute the following amended paragraph:

The conditions stated above, permit the formation of an unprotected oligonucleotide which is equipped with the activated thioester. Subsequent reaction with a second peptide containing a terminal cysteine residue, permits a facile coupling with the formation of a native peptide bond and can generate oligopeptide chains of 100 or more amino acid residues (FIG. 1).

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Page 21, line 26 after last paragraph, please insert the following ten paragraphs directed to the Brief Description of Drawings Section, viz.:

Brief Description of Drawings:

Figure 1 illustrates the overall principle of 'native chemical ligation.'

Figure 2 illustrates a general synthesis of Boc-amino thioester linker (1), dicyclohexylamine salt.

Figure 3 illustrates the progress of a first native chemical ligation reaction for the production of model peptide (Sequence No. 5) of Example 1 using a first peptide segment (Sequence No. 2) and a second peptide segment (Sequence No. 3) having a S-benzyl derivative, as monitored by HPLC.

Figure 4 illustrates the progress of a second native chemical ligation reaction for the production of model peptide (Sequence No. 5) of Example 1 using a first peptide segment (Sequence No. 2) and a second peptide segment (Sequence No. 6) having a thioacetic acid derivative, as monitored by HPLC.

Figure 5 illustrates the kinetics of the native chemical ligation reaction of Example 2 with respect to the linkage of peptide segments 46-76 and 77-95 for producing IL-3 receptor  $\beta$ -subunit.

Figure 6 illustrates the reversed-phase HPLC purification and electrospray mass spectroscopy of unreacted IL-3 peptide segments (46-76) and (77-95) of IL-3 receptor  $\beta$ -subunit and of their native chemical ligation product as detailed in Example 2.

Figure 7 illustrates a native chemical ligation scheme for producing LI-8 as detailed in Example 3.

Figure 8 illustrates the reversed-phase HPLC purification and electrospray mass spectroscopy of IL-8 (Sequence No.: 10) as formed by a native chemical ligation of peptides Sequence No.'s 8 and 9 in Example 3. Figure 8B illustrates the unreacted IL-8 peptides (1-33)-SBenzyl (Sequence No. 8) and IL-8 34-72 (Sequence No. 9) prior to ligation;

Figure 8C illustrates the unfolded IL-8 ligation product, Sequence No. 10; and Figure 8D illustrates the folded IL-8 ligation product, Sequence No. 10.

Figure 9 illustrates the reversed-phase HPLC purification and electrospray mass spectroscopy of HIV-1 K41 protease (Sequence No.: 15) as formed by native chemical ligation of peptides Sequence No.'s 11 and 12 in Example 4.

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Figure 10 illustrates the reversed-phase HPLC purification of Barnase K39 (Sequence No.: 20) as formed by native chemical ligation of peptide Sequence No's. 17 and 18 in Example 5.

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Page 29, please delete the first full paragraph, lines 2-14, and substitute the following amended paragraph:

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Another, unpublished model using the 2-thioacetic acid derivative Leu-Tyr-Arg-Ala-Gly-SCH<sub>2</sub>COOH (SEQ ID NO 6), formed from attack of the thioacid Leu-Tyr-Arg-Ala-Gly-SH (SEQ ID NO 1), onto 2-bromoacetic acid in methylene chloride) + Cys-Arg-Ala-Glu-Tyr-Ser (Sequence No.: 2) was ligated at pH 6.8 in 0.2 M phosphate buffer, at 45 °C. After 1.0 hour the reaction had proceeded to 80% as observed in FIG. 4 by HPLC. The isolation of oxidation products from the ligated Leu-Tyr-Arg-Ala-Gly-Cys-Arg-Ala-Glu-Tyr-Ser (SEQ ID NO 5) and unreacted Cys-Arg-Ala-Glu-Tyr-Ser (SEQ ID NO 2) demonstrated the presence of a free thiol ligation product.

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In the Sequence Listing:

Please delete the originally-filed paper copy of the sequence listing and substitute the paper copy of the sequence listing submitted on August 1, 2001.

Sequence Listing:

Please delete the originally-filed paper copy of the sequence listing and substitute the paper copy of the corrected sequence listing submitted on August 1, 2001.

The sequence listings for SEQ ID NO:4 and SEQ ID NO:11 contained in the Sequence Listing filed on August 1, 2001 differ from the corresponding sequences contained in the originally filed Sequence Listing.

SEQ ID NO 4 has been corrected by clarifying the modification of the glycine at position 5. Support for this correction is found in the Specification at page 28, lines 11-13.

SEQ ID NO 11 has been corrected by deleting the reference to an Xaa at position 27. The sequence itself correctly indicates that there is a glycine at position 27. Support for this correction, i.e., deletion of a reference to an Xaa at position 27, is found in the sequence itself and in the Specification at page 37, lines 19-20.

Certification:

I hereby state that the content of the paper and computer readable copies of the Sequence Listing, submitted in accordance with 37 CFR §1.821 - §1.825, are the same, except where noted above.

I hereby state that the Amendments, made in accordance with 37 CFR §1.821 - §1.825 do not include new matter.

I hereby declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and